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Last updated by author(s):	Jan 21, 2021

Reporting Summary

Nature Research wishes to improve the reproducibility of the work that we publish. This form provides structure for consistency and transparency in reporting. For further information on Nature Research policies, see our <u>Editorial Policies</u> and the <u>Editorial Policy Checklist</u>.

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For	r all statistical analyses, (confirm that the following items are present in the figure legend, table legend, main text, or Methods section.
n/a	a Confirmed	
	The exact sample	size (n) for each experimental group/condition, given as a discrete number and unit of measurement
	A statement on w	hether measurements were taken from distinct samples or whether the same sample was measured repeatedly
	The statistical test Only common tests	t(s) used AND whether they are one- or two-sided should be described solely by name; describe more complex techniques in the Methods section.
	A description of a	Il covariates tested
	A description of a	ny assumptions or corrections, such as tests of normality and adjustment for multiple comparisons
	A full description AND variation (e.	of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)
	For null hypothes Give P values as exa	is testing, the test statistic (e.g. F , t , r) with confidence intervals, effect sizes, degrees of freedom and P value noted ct values whenever suitable.
\boxtimes	For Bayesian anal	ysis, information on the choice of priors and Markov chain Monte Carlo settings
X	For hierarchical a	nd complex designs, identification of the appropriate level for tests and full reporting of outcomes
	Estimates of effec	t sizes (e.g. Cohen's d , Pearson's r), indicating how they were calculated
	•	Our web collection on statistics for biologists contains articles on many of the points above

Software and code

Policy information about availability of computer code

Data collection

STED images were collected using a Leica TCS SP8 gated STED (gSTED) 3X superresolution system (Leica Microsystems) equipped with a tunable white light laser, CW 592 nm and 660 nm depletion lines, and a pulsed 775 nm depletion line. Live cell images were collected with either Leica SP8 or confocal spinning disk with a Yokogawa CSU-10 and a Hamamatsu EM-CCD digital camera attached to an inverted Leica microscope and controlled by Metamorph software 7.10.

Data analysis

No programs were written for this manuscript. Image analysis was conducted off-line using Fiji Image J (https://imagej.net/Fiji) and built-in macros. 3D rendering of STED images was performed using the Imaris software (Bitplane AG 8.3.1). Statistics was performed using Graphpad Prism 8.0. Power calculations were performed using G*Power software 3.1.

For manuscripts utilizing custom algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors and reviewers. We strongly encourage code deposition in a community repository (e.g. GitHub). See the Nature Research guidelines for submitting code & software for further information.

Data

Policy information about availability of data

 $All\ manuscripts\ must\ include\ a\ \underline{data\ availability\ statement}.\ This\ statement\ should\ provide\ the\ following\ information,\ where\ applicable:$

- Accession codes, unique identifiers, or web links for publicly available datasets
- A list of figures that have associated raw data
- A description of any restrictions on data availability

The data that support the findings of this study are available from the corresponding author upon reasonable request. There are no restriction on data availability. The data availability statement is included with the manuscript.

Field-spe	ecific reporting				
\times Life sciences	ne below that is the best fit for your research. If you are not sure, read the appropriate sections before making your selection. Behavioural & social sciences Ecological, evolutionary & environmental sciences che document with all sections, see nature.com/documents/nr-reporting-summary-flat.pdf				
Life scier	nces study design				
All studies must dis	close on these points even when the disclosure is negative.				
Sample size	ndard power calculations using the power of 0.8-0.95 and alpha = 0.05 were used to determine the sample size. Power calculations were formed in the G*power 3.1 software using tests and variance based on our previously published data (Hruska et al., 2015, Hruska et al., 18, Dalva et al., 2000, Kayser et al., 2008, McClelland et al., 2010 and Nolt et al., 2011).				
Data exclusions	No data were excluded from our analyses.				
Replication	Each experiment was replicated a minimum of three times and data was reliably reproduced with each replication attempt.				
Randomization	a were acquired and analyzed based on the standards in the field and our previously published results (Hruska et al., 2015, Hruska et al., 8, Dalva et al., 2000, Kayser et al., 2008, McClelland et al., 2010 and Nolt et al., 2011). However, no method of randomization was used to ermine how samples were allocated to experimental groups and processed since the experiments were done entirely in vitro. Because no nals were used in ayn experimental groups, randomization was not relevant.				
Blinding	Data for spine analysis of module number represent observations and were acquired and analyzed with an experimenter blinded to the condition. Experimenter as well as analyzer were blinded to conditions.				
We require informati	g for specific materials, systems and methods on from authors about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material ted is relevant to your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.				
Materials & ex	perimental systems Methods				
n/a Involved in th	ne study n/a Involved in the study				
Antibodies					
Eukaryotic					
	☐ Palaeontology and archaeology ☐ MRI-based neuroimaging ☐ Animals and other organisms				
Clinical data					
	Dual use research of concern				
Antibodies					
Antibodies used	Primary antibodies: mouse monoclonal (IgG2A) anti-PSD-95 clone K28/43(1:200, Neuromab, UC Davis, Davis, CA), mouse monoclonal IgG1 anti-PSD-95 clone 7E3-1B8 (1:250, Thermo Fisher Scientific, Waltham, MA, cat#: MA1-046) anti-guinea pig polyclonal anti-Bassoon (1:300, Synaptic Systems, Gottingen, Germany, cat# 141004), rabbit polyclonal anti-Bassoon (1:300, Synaptic Systems, cat #: 141 003), chicken anti-GFP (1:2000, Abcam, Cambridge, MA, cat# ab13970), rabbit monoclonal anti-GluN1 (1:500, AB9864, Millipore Sigma), mouse monoclonal (IgG2A) anti-GluN2A clone N327/95 (1:250, Neuromab), mouse monoclonal (IgG2B) anti-GluN2B clone N59/3625(1:250, Neuromab), mouse monoclonal (IgG1) anti-GluA1 clone N355/11 (1:250, Neuromab), rabbit polyclonal anti-GluA2 (1:500, Synaptic Systems, cat#: 182 103),				

Secondary antibodies: Goat anti mouse IgG2A Atto 425 (1:250, Rockland, Inc.,cat# 610-151-041), Goat anti-mouse IgG1 Atto-425 (1:250, Rockland, cat#: 610-151-040),

mouse monoclonal (IgG2A) anti-Synaptotagmin 1 (1:500, Synaptic Systems, cat#: 105011);

Goat anti-rabbit Atto 425 (1:250, Rockland, cat#: 611-151-122),

Goat anti-mouse IgG1 Atto-647N (1:500, Rockland, Inc., cat # 610-156-040),
Goat anti-mouse IgG2A Atto-647N (1:500, Rockland, cat#: 610-156-041),
Goat anti-mouse IgG2B Atto-647N (1:500, Rockland, cat#: 610-156-042),
Goat anti-rabbit Atto-647N (1:500, Rockland, Inc., cat # 611-156-122),
Goat anti-mouse IgG1 AlexaFluor-594 (1:500, Jackson ImmunoResearch, cat# 115-587-185),
Goat anti-mouse IgG2A AlexaFluor-594 (1:500, Jackson ImmunoResearch, cat# 115-585-206),
Donkey anti guinea pig AlexaFluor-594 (1:500, Jackson ImmunoResearch, cat # 706-586-148)
Donkey anti-rabbit AlexaFluor-594 (1:500, Jackson ImmunoResearch, cat # 711-585-152).

Validation

All primary antibodies were validated in the KO animals by the respective vendors. Additionally, all primary antibodies were profiled in previous publications and were reported to be specific:

mouse monoclonal (IgG2A) anti-PSD-95 clone K28/43(1:200, Neuromab, UC Davis, Davis, CA) (Hruska et al., 2015; Hruska et al., 2018) mouse monoclonal IgG1 anti-PSD-95 clone 7E3-1B8 (1:250, Thermo Fisher Scientific, Waltham, MA, cat#: MA1-046) (Schaffer et al., 2018; Abrahamsson et al., 2017)

anti-guinea pig polyclonal anti-Bassoon (1:300, Synaptic Systems, Gottingen, Germany, cat# 141004) (Hruska et al., 2018; Schroeder et al., 2021)

rabbit polyclonal anti-Bassoon (1:300, Synaptic Systems, cat #: 141 003) (Hruska et al., 2018; Nishida et al., 2021) chicken anti-GFP (1:2000, Abcam, Cambridge, MA, cat# ab13970) (Greer et al., 2021; Steevens et al., 2021) rabbit monoclonal anti-GluN1 (1:500, AB9864, Millipore Sigma) (Gleichman et al., 2012; Peng et al., 2014) mouse monoclonal (IgG2A) anti-GluN2A clone N327/95 (1:250, Neuromab) (Balsara et al., 2015) mouse monoclonal (IgG2B) anti-GluN2B clone N59/3625(1:250, Neuromab) (Washburn et al., 2020; Hruska et al., 2015) mouse monoclonal (IgG1) anti-GluA1 clone N355/11 (1:250, Neuromab) (Moraga-Amaro et al., 2016) rabbit polyclonal anti-GluA2 (1:500, Synaptic Systems, cat#: 182 103) (Won et al., 2019; Fiuza et al., 2017) mouse monoclonal (IgG2A) anti-Synaptotagmin 1 (1:500, Synaptic Systems, cat#: 105011) (Farsi et al., 2021; Hoffmann-Conaway et al., 2020)

Animals and other organisms

Policy information about studies involving animals; ARRIVE guidelines recommended for reporting animal research

Laboratory animals

Long-Evans E17-18 rat embryos from timed pregnant animals purchased from Charles River Laboratories Inc. (Wilmington, MA) were used to make primary cortical neuron cultures. Both male and female embryos were used.

Wild animals

The study did not involve the use of wild animals.

Field-collected samples

The study did not involve samples collected from the field.

Ethics oversight

All animal studies were approved by the Institutional Animal Care and Use Committee guidelines at Thomas Jefferson University in accordance with the US National Institutes of Health guidelines.

Note that full information on the approval of the study protocol must also be provided in the manuscript.